



#4

1

SEQUENCE LISTING

<110> SCHLAGER, JOHN J.
SWEENEY, RICHARD E.
AVERY, DOUGLAS P.

<120> AUTOMATED METHOD OF IDENTIFYING AND ARCHIVING NUCLEIC
ACID SEQUENCES

<130> RICD-00-21

<140> 09/961,058
<141> 2001-09-24

<150> 60/235,899
<151> 2000-09-28

<160> 16

<170> PatentIn Ver. 2.1

<210> 1
<211> 404
<212> DNA
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Comparative DNA
sequence

<220>
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<222> (71)
<223> a, t, c or g

<220>
<221> modified_base
<222> (142)
<223> a, t, c or g

<220>
<221> modified_base
<222> (224)..(225)
<223> a, t, c or g

<220>
<221> modified_base
<222> (237)
<223> a, t, c or g

<220>
<221> modified_base
<222> (289)
<223> a, t, c or g

<220>
<221> modified_base
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<223> a, t, c or g

<220>

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<222> (301)

<223> a, t, c or g

<220>

<221> modified_base

<222> (323)

<223> a, t, c or g

<220>

<221> modified_base

<222> (333)

<223> a, t, c or g

<220>

<221> modified_base

<222> (350)

<223> a, t, c or g

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<222> (357)

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acaatggcctt ttcagttcct anaggacaca ttgtgagcaa tctcagcaca gtaagatttg 180
ttgcacatca gcagcacctc cagctccttg acattgtgga ccannaactt gcggaanccg 240
ctgggcagca tgtgcttggt tttcttggtg ctcccacaac cgaagttnng gcatcangat 300
ntggcccttg aaccttctcc ccncctggtg tcnatgcctc tgggtttccn catttcnctt 360
aatttcccat atcgggtctga cttaattttc acatatcggt ctga 404
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<210> 2

<211> 383

<212> DNA

<213> Mus musculus

<400> 2

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tcgctgcgta gcctggcggt gggattgggt actctgatgg ccagctgtgc tgctctttct 120
acaatggcctt ttcggttctt agaggacaca ttgtgagcaa tctcagcaca gtaagatttg 180
ttgcacatca gcagcacctc cagctccttg acattgtgga ccaggaactt gcggaagccg 240
ctgggcagca tgtgcttggt tttcttggtg ctcccataac cgatgttggg catcaggatc 300
tggcccttga accttctccg caccctggtg tcaatgcctc tgggtttccg ccagtttcgc 360
ttaattttca catatcggtc tga 383
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<210> 3

<211> 349

<212> DNA

<213> Unknown Organism

<220>
 <223> Description of Unknown Organism: Comparative DNA
 sequence

<220>
 <221> modified_base
 <222> (71)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (142)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (224)..(225)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (237)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (289)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (297)
 <223> a, t, c or g

<220>
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 <222> (301)
 <223> a, t, c or g

<220>
 <221> modified_base
 <222> (323)
 <223> a, t, c or g

<220>
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 <222> (333)
 <223> a, t, c or g

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 acaatggcctt ttcagttctt anaggacaca ttgtgagcaa tctcagcaca gtaagatttg 180
 ttgcacatca gcagcacctc cagctccttg acattgtgga ccannaactt gcggaanccg 240
 ctgggcagca tgtgcttggg tttcttggtg ctcccacaac cgaagtttng gcatcangat 300
 ntggcccttg aaccttctcc ccnctgttg tcnatgcctc tgggtttcc 349

<210> 4
 <211> 349
 <212> DNA
 <213> Rattus sp.

<400> 4
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 acgatggcct ttcggttcct agaggacaca ttgtgagcaa tctcagcaca gtaagatttg 180
 ttgcacatca gcagcacttc cagctccttg acattgtgga ccagaaactt ccggaagccg 240
 ctaggcagca tgtgcttggt tttcttgta ctcccgtaac caatgttggg catcaggatc 300
 tggcccttga atcttctccg caccctgttg tcgatgcctc tgggtttcc 349

<210> 5
 <211> 103
 <212> DNA
 <213> Unknown Organism

<220>
 <223> Description of Unknown Organism: Illustrative DNA
 sequence

<400> 5
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 acataaaaac tgcacacaag ccatctactc attttcttcg ctg 103

<210> 6
 <211> 20
 <212> DNA
 <213> Unknown Organism

<220>
 <223> Description of Unknown Organism: Adapter sequence

<400> 6
 agcggccgcc cgggcaggtc 20

<210> 7
 <211> 20
 <212> DNA
 <213> Unknown Organism

<220>
 <223> Description of Unknown Organism: Adapter sequence

<400> 7
 acctcggccg cgaccacgct 20

<210> 8
 <211> 33
 <212> DNA
 <213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 8

ttactagtgg atccgagctc ggtaccaagc ttc

33

<210> 9

<211> 20

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 9

agcgtggtcg cggccgaggt

20

<210> 10

<211> 20

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 10

acctgcccgg gcggccgctc

20

<210> 11

<211> 33

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 11

cacactggcg gccgctcgag catgcatcta gag

33

<210> 12

<211> 19

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 12

agcggccgcc cgggcaggt

19

<210> 13

<211> 19

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Adapter sequence

<400> 13

acctgcccgg gcggccgct

19

<210> 14

<211> 624

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Illustrative DNA
sequence

<220>

<221> modified_base

<222> (372)

<223> a, t, c or g

<220>

<221> modified_base

<222> (403)

<223> a, t, c or g

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<221> modified_base

<222> (407)

<223> a, t, c or g

<220>

<221> modified_base

<222> (434)

<223> a, t, c or g

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<221> modified_base

<222> (436)

<223> a, t, c or g

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<221> modified_base

<222> (453)

<223> a, t, c or g

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<221> modified_base

<222> (483)

<223> a, t, c or g

<220>

<221> modified_base

<222> (492)

<223> a, t, c or g

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 <222> (500)
 <223> a, t, c or g

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 <223> a, t, c or g

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<220>
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 <222> (546)
 <223> a, t, c or g

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 <222> (550)
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 <222> (560)
 <223> a, t, c or g

<220>
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 <222> (570)
 <223> a, t, c or g

<220>
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 <222> (575)
 <223> a, t, c or g

<220>
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 <222> (613)
 <223> a, t, c or g

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 agaaagacct agaaggttgt agatgggaaa tcaggaatga tttgaactga taaagatttc 120
 ggactcataa gaacacattt tataaatgtt aaacacaaaa actacatgac tgaagataga 180
 agagaatgcg atggatttta ttacacatgg tggagagag aagaggcgtg taggtttgca 240
 aacaaagtta agaaatagga aactgaattt ttcattgtac agaaaatgta tctcttgggg 300
 aaggcctgtg tacctgcccg ggcggccgct cgaaattcca gcacactggc ggccgttact 360
 agtggatccc anctcgggtac caagcttggt gttatcatgg tcntaanctg tttcctgtgt 420

```

gaaattgtta tccncccc attcccccc acnttccaac ccgaaacett aaatttttaa 480
ccnggggtgc cnaatgaatn acccaccnccn ttattgcttt gccnccctgcc ccttcctcg 540
gaaccntctn cccctctttn taaaccgcn cccnggaaa gcgtttcttt tggccctcc 600
cctccccctc ctnatcctgc ccct                                     624

```

```

<210> 15
<211> 32
<212> DNA
<213> Artificial Sequence

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```

<220>
<223> Description of Artificial Sequence: Primer

```

```

<400> 15
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```

```

<210> 16
<211> 20
<212> DNA
<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Primer

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<400> 16
ttcgaaccat ggctcgagcc                                     20

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